

Exhibit E

<!--StartFragment-->RESULT 2

ABT07740

ID ABT07740 standard; DNA; 2119 BP.

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AC ABT07740;

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DT 14-NOV-2002 (first entry)

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DE Breast cancer-associated gene sequence 48.

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KW Gene; ds; breast cancer; breast cancer-associated gene sequence;

KW drug development; pharmacogenetics; biosensor development.

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OS Unidentified.

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PN WO200259377-A2.

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PD 01-AUG-2002.

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PF 24-JAN-2002; 2002WO-US002242.

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PR 24-JAN-2001; 2001US-0263965P.

PR 02-FEB-2001; 2001US-0265928P.

PR 09-APR-2001; 2001US-00829472.

PR 09-APR-2001; 2001US-0282698P.

PR 04-MAY-2001; 2001US-0288590P.

PR 29-MAY-2001; 2001US-0294443P.

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PA (EOSB-) EOS BIOTECHNOLOGY INC.

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PI Mack DH, Gish KC, Afar D;

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DR WPI; 2002-583738/62.

DR N-PSDB; ABJ05583.

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PT Detecting a breast cancer-associated transcript in a patient's cell,
 PT useful for diagnosing breast cancer, comprises contacting a biological
 PT sample with a polynucleotide that selectively hybridizes with breast
 PT cancer nucleic acids.

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PS Claim 9; Page 388-389; 414pp; English.

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CC The invention comprises a method of detecting a breast cancer-associated
 CC transcript in a cell from a patient. The method of the invention involves
 CC contacting a biological sample from the patient with a nucleotide that
 CC hybridises to one of the 69 breast cancer-associated gene sequences shown
 CC in the specification. The method of the invention is useful in the
 CC diagnosis or prognosis of breast cancer, and for detecting genes that are
 CC up or down-regulated in breast cancer cells. Genes identified by the
 CC method of the invention can be used in diagnostic purposes and also as
 CC targets for screening for therapeutic compounds that modulate breast
 CC cancer (e.g. hormones or antibodies). Identification of genes that are
 CC over or under expressed in breast cancer can additionally provide high-
 CC resolution, high-sensitivity datasets which can be used in the areas of
 CC diagnostics, therapeutics, drug development, pharmacogenetics, protein
 CC structure and biosensor development. DNA sequences ABT07693 - ABT07761
 CC represent the 69 breast cancer-associated gene sequences of the invention

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SQ Sequence 2119 BP; 646 A; 389 C; 492 G; 592 T; 0 U; 0 Other;

Query Match 100.0%; Score 2119; DB 6; Length 2119;

Best Local Similarity 100.0%; Pred. No. 0;

Matches 2119; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy	1	GGCACGAGTAGGGGTGGCGGGTCAGTGTGCTCGGGGGCTTCTCCATCCAGGTCCCTGGA	60
Db	1	GGCACGAGTAGGGGTGGCGGGTCAGTGTGCTCGGGGGCTTCTCCATCCAGGTCCCTGGA	60
Qy	61	GTTCTGGTCCCTGGAGCTCCGCACTTGGCGCGCAACCTGCGTGAGGCAGCGGACTCTG	120
Db	61	GTTCTGGTCCCTGGAGCTCCGCACTTGGCGCGCAACCTGCGTGAGGCAGCGGACTCTG	120
Qy	121	GCGACTGGCCGGCCATGCCTTCCCGGGCTGAGGACTATGAAGTGTGTACACCATTGGCA	180
Db	121	GCGACTGGCCGGCCATGCCTTCCCGGGCTGAGGACTATGAAGTGTGTACACCATTGGCA	180

Qy	181	CAGGCTCCTACGGCCGCTGCCAGAAGATCCGGAGGAAGAGTGATGGCAAGATATTAGTTT	240
Db	181	CAGGCTCCTACGGCCGCTGCCAGAAGATCCGGAGGAAGAGTGATGGCAAGATATTAGTTT	240
Qy	241	GGAAAGAACTTGACTATGGCTCCATGACAGAAGCTGAGAAACAGATGCTTGTTCCTGAAG	300
Db	241	GGAAAGAACTTGACTATGGCTCCATGACAGAAGCTGAGAAACAGATGCTTGTTCCTGAAG	300
Qy	301	TGAATTGCTTCGTGAACCTGAAACATCCAAACATCGTTCGTTACTATGATCGGATTATTG	360
Db	301	TGAATTGCTTCGTGAACCTGAAACATCCAAACATCGTTCGTTACTATGATCGGATTATTG	360
Qy	361	ACCGGACCAATACAACACTGTACATTGTAATGGAATATTGTGAAGGAGGGGATCTGGCTA	420
Db	361	ACCGGACCAATACAACACTGTACATTGTAATGGAATATTGTGAAGGAGGGGATCTGGCTA	420
Qy	421	GTGTAATTACAAAGGGAACCAAGGAAAGGCAATACTTAGATGAAGAGTTTGTTCCTCGAG	480
Db	421	GTGTAATTACAAAGGGAACCAAGGAAAGGCAATACTTAGATGAAGAGTTTGTTCCTCGAG	480
Qy	481	TGATGACTCAGTTGACTCTGGCCCTGAAGGAATGCCACAGACGAAGTGATGGTGGTCATA	540
Db	481	TGATGACTCAGTTGACTCTGGCCCTGAAGGAATGCCACAGACGAAGTGATGGTGGTCATA	540
Qy	541	CCGTATTGCATCGGGATCTTAAACCAGCCAATGTTTTCTGGATGGCAAGCAAAACGTCA	600
Db	541	CCGTATTGCATCGGGATCTTAAACCAGCCAATGTTTTCTGGATGGCAAGCAAAACGTCA	600
Qy	601	AGCTTGGAGACTTTGGGCTAGCTAGAATATTAAACCATGACACGAGTTTGC AAAACAT	660
Db	601	AGCTTGGAGACTTTGGGCTAGCTAGAATATTAAACCATGACACGAGTTTGC AAAACAT	660
Qy	661	TTGTTGGCACACCTTATTACATGTCTCCTGAACAAATGAATCGCATGTCTACAATGAGA	720
Db	661	TTGTTGGCACACCTTATTACATGTCTCCTGAACAAATGAATCGCATGTCTACAATGAGA	720
Qy	721	AATCAGATATCTGGTCATTGGGCTGCTTGCTGTATGAGTTATGTGCATTAATGCCTCCAT	780
Db	721	AATCAGATATCTGGTCATTGGGCTGCTTGCTGTATGAGTTATGTGCATTAATGCCTCCAT	780
Qy	781	TTACAGCTTTTAGCCAGAAAGAAGCTCGCTGGGAAAATCAGAGAAGGCAAATTCAGGCGAA	840
Db	781	TTACAGCTTTTAGCCAGAAAGAAGCTCGCTGGGAAAATCAGAGAAGGCAAATTCAGGCGAA	840
Qy	841	TTCCATACCGTTACTCTGATGAATTGAATGAAATTATTACGAGGATGTAAACTTAAAGG	900
Db	841	TTCCATACCGTTACTCTGATGAATTGAATGAAATTATTACGAGGATGTAAACTTAAAGG	900
Qy	901	ATTACCATCGACCTTCTGTTGAAGAAATTCCTTGAGAACCCCTTAATAGCAGATTTGGTTG	960
Db	901	ATTACCATCGACCTTCTGTTGAAGAAATTCCTTGAGAACCCCTTAATAGCAGATTTGGTTG	960
Qy	961	CAGACGAGCAAAAGAAGAAATCTTGAGAGAAGAGGGCGACAATTAGGAGAGCCAGAAAAAT	1020
Db	961	CAGACGAGCAAAAGAAGAAATCTTGAGAGAAGAGGGCGACAATTAGGAGAGCCAGAAAAAT	1020
Qy	1021	CGCAGGATTCCAGCCCTGTATTGAGTGAGCTGAAACTGAAGGAAATTCAGTTACAGGAGC	1080
Db	1021	CGCAGGATTCCAGCCCTGTATTGAGTGAGCTGAAACTGAAGGAAATTCAGTTACAGGAGC	1080
Qy	1081	GAGAGCGAGCTCTCAAAGCAAGAGAAGAAAGATTGGAGCAGAAAGAACAGGAGCTTTGTG	1140
Db	1081	GAGAGCGAGCTCTCAAAGCAAGAGAAGAAAGATTGGAGCAGAAAGAACAGGAGCTTTGTG	1140
Qy	1141	TTCGTGAGAGACTAGCAGAGGACAAACTGGCTAGAGCAGAAAAATCTGTTGAAGAACTACA	1200
Db	1141	TTCGTGAGAGACTAGCAGAGGACAAACTGGCTAGAGCAGAAAAATCTGTTGAAGAACTACA	1200
Qy	1201	GCTTGCTAAAGGAACGGAAGTTCCTGTCTCTGGCAAGTAATCCAGAACTTCTTAATCTTC	1260
Db	1201	GCTTGCTAAAGGAACGGAAGTTCCTGTCTCTGGCAAGTAATCCAGAACTTCTTAATCTTC	1260
Qy	1261	CATCCTCAGTAATTAAGAAGAAAGTTTCAATTCAGTGGGAAAGTAAGAGAACATCATGA	1320
Db	1261	CATCCTCAGTAATTAAGAAGAAAGTTTCAATTCAGTGGGAAAGTAAGAGAACATCATGA	1320

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Qy      1321 GGAGTGAGAAATTCTGAGAGTCAGCTCACATCTAAGTCCAAGTGCAAGGACCTGAAGAAAA 1380
        |||
Db      1321 GGAGTGAGAAATTCTGAGAGTCAGCTCACATCTAAGTCCAAGTGCAAGGACCTGAAGAAAA 1380

Qy      1381 GGCTTCACGCTGCCCAGCTGCGGGCTCAAGCCCTGTCAGATATTGAGAAAAATTACCAAC 1440
        |||
Db      1381 GGCTTCACGCTGCCCAGCTGCGGGCTCAAGCCCTGTCAGATATTGAGAAAAATTACCAAC 1440

Qy      1441 TGAAAAGCAGACAGATCCTGGGCATGCGCTAGCCAGGTAGAGAGACACAGAGCTGTGTAC 1500
        |||
Db      1441 TGAAAAGCAGACAGATCCTGGGCATGCGCTAGCCAGGTAGAGAGACACAGAGCTGTGTAC 1500

Qy      1501 AGGATGTAATATTACCAACCTTTAAAGACTGATATTCAAATGCTGTAGTGTGAATACTT 1560
        |||
Db      1501 AGGATGTAATATTACCAACCTTTAAAGACTGATATTCAAATGCTGTAGTGTGAATACTT 1560

Qy      1561 GGCCCCATGAGCCATGCCTTTCTGTATAGTACACATGATATTTCGGAATTGGTTTTACTG 1620
        |||
Db      1561 GGCCCCATGAGCCATGCCTTTCTGTATAGTACACATGATATTTCGGAATTGGTTTTACTG 1620

Qy      1621 TTCTTCAGCAACTATTGTACAAAATGTTACATTTAATTTTCTTTCTTTTAAAGAAC 1680
        |||
Db      1621 TTCTTCAGCAACTATTGTACAAAATGTTACATTTAATTTTCTTTCTTTTAAAGAAC 1680

Qy      1681 ATATTATAAAAAGAATACTTTCTTGTTGGGCTTTTAAATCCTGTGTGTGATTACTAGTAG 1740
        |||
Db      1681 ATATTATAAAAAGAATACTTTCTTGTTGGGCTTTTAAATCCTGTGTGTGATTACTAGTAG 1740

Qy      1741 GAACATGAGATGTGACATTCTAAATCTTGGGAGAAAAAATAATATTAGGAAAAAATATT 1800
        |||
Db      1741 GAACATGAGATGTGACATTCTAAATCTTGGGAGAAAAAATAATATTAGGAAAAAATATT 1800

Qy      1801 TATGCAGGAAGAGTAGCACTCACTGAATAGTTTTAAATGACTGAGTGGTATGCTTACAAT 1860
        |||
Db      1801 TATGCAGGAAGAGTAGCACTCACTGAATAGTTTTAAATGACTGAGTGGTATGCTTACAAT 1860

Qy      1861 TGTCTGTCTAGATTTAAATTTTAAAGTCTGAGATTTTAAATGTTTTGAGCTTAGAAAAC 1920
        |||
Db      1861 TGTCTGTCTAGATTTAAATTTTAAAGTCTGAGATTTTAAATGTTTTGAGCTTAGAAAAC 1920

Qy      1921 CCAGTTAGATGCAATTTGGTCATTAATACCATGACATCTTGCTTATAAATATTCCATTGC 1980
        |||
Db      1921 CCAGTTAGATGCAATTTGGTCATTAATACCATGACATCTTGCTTATAAATATTCCATTGC 1980

Qy      1981 TCTGTAGTTCAAATCTGTTAGCTTTGTGAAAATTCATCACTGTGATGTTTGTATTCTTTT 2040
        |||
Db      1981 TCTGTAGTTCAAATCTGTTAGCTTTGTGAAAATTCATCACTGTGATGTTTGTATTCTTTT 2040

Qy      2041 TTTTTTCTGTTTAAACAGAAATATGAGCTGTCTGTCATTACCTACTTCTTTCCCACTAAA 2100
        |||
Db      2041 TTTTTTCTGTTTAAACAGAAATATGAGCTGTCTGTCATTACCTACTTCTTTCCCACTAAA 2100

Qy      2101 TAAAAGAATTCTTCAGTTA 2119
        |||
Db      2101 TAAAAGAATTCTTCAGTTA 2119
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